03AD

OIPE

#7

RAW SEQUENCE LISTING   DATE: 05/13/2002 P.6
PATENT APPLICATION: US/09/667,170A   TIME: 15:12:12

Input Set : A:\475c7.app
Output Set: N:\CRF3\05132002\I667170A.raw

```
  3 <110> APPLICANT: Reed, Steven G.
  4         Henderson, Robert A.
  5         Lodes, Michael J.
  6         Fling, Steven P.
  7         Mohamath, Raodoh
  8         Algate, Paul A.
  9         Secrist, Heather
 10         Indirias, Carol Yoseph
 11         Benson, Darin R.
 13 <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
 14         THE THERAPY AND DIAGNOSIS OF LUNG CANCER
 17 <130> FILE REFERENCE: 210121.475C7
 19 <140> CURRENT APPLICATION NUMBER: US 09/667,170A
C--> 20 <141> CURRENT FILING DATE: 2002-05-03
 22 <160> NUMBER OF SEQ ID NOS: 440
 24 <170> SOFTWARE: FastSEQ for Windows Version 3.0
 26 <210> SEQ ID NO: 1
 27 <211> LENGTH: 339
 28 <212> TYPE: DNA
 29 <213> ORGANISM: Homo sapien
 31 <400> SEQUENCE: 1
 32  gtactcagac aggatagtca tcatgtagca caaagcamat cctgtttcta tacttgtagt      60
 33  ttgctctcac tcagtggcat ratcattact atacagtgta gaatgttrtt atgtagcata     120
 34  gatgtggggt ctctagccca cagctctsta cctttgtcta gcactcctgt cctcatacct     180
 35  ragtggcctg tccatcagca tgtttctcat ctactttgct tgtccagtcc actgtggtcc     240
 36  tcccttgccc tctcccttat gtggcagagt ggaaccagct gtcctgagac ttgagttcaa     300
 37  catctggttc gcccatytgc atgtttgtgg tctgagtac                           339
 39 <210> SEQ ID NO: 2
 40 <211> LENGTH: 698
 41 <212> TYPE: DNA
 42 <213> ORGANISM: Homo sapien
 44 <220> FEATURE:
 45 <221> NAME/KEY: misc_feature
 46 <222> LOCATION: (1)...(698)
 47 <223> OTHER INFORMATION: n = A,T,C or G
 49 <400> SEQUENCE: 2
 50  gtactcagac cacgactgca ttttctccac tgctgacggg tctaatacca gctgcttccc      60
W--> 51  tttcttggag gcagagctng tgaccttgag aaagtgacct gtgaccatca tgtgggtagt     120
 52  gagctgctgc aaggtgtcat gggagctccc acactccatg cactttwaga tctgggactt     180
 53  gcaggcctca ractgccagg tgtagctcgc tccattttgg tagccatagc gsttgttgga     240
 54  ggacaactgc aagttggcgt tcttctgaga agaaaaagaa tctgcaaaag atcctgtggt     300
 55  tgaatcgggg gaacacggcc gattgacatc aaaaacgcgt ttcttagccc gggtgaccat     360
 56  tttcgaggaa atggttgggg actggctcct tcaaaggcac ttttttggtta tgttttgttt     420
```

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                      RAW SEQUENCE LISTING            DATE: 05/13/2002
                      PATENT APPLICATION:  US/09/667,170A    TIME: 15:12:12

                      Input Set : A:\475c7.app
                      Output Set: N:\CRF3\05132002\I667170A.raw

W--> 57  yaatcatgtk gacgctccaa tcttggragg gaatcgaang rantcnccnc caaaacatrc      480
     58  stttcagraa ccttttgarc atcctctttt ttccgtrtcc cggmaargcc cytttccckg      540
W--> 59  ggctttgaaa wyagcctsgt tgggttctta aattaccart ccacnwgttg gaattccccg      600
W--> 60  ggcccctgc ccggktccaa ccaattttgg graaaacccc cncansccgt tkggantgcn       660
W--> 61  acaacntggn nttttttcntt tcgtgntccc ctngaacc                            698
     63  <210> SEQ ID NO: 3
     64  <211> LENGTH: 697
     65  <212> TYPE: DNA
     66  <213> ORGANISM: Homo sapien
     68  <220> FEATURE:
     69  <221> NAME/KEY: misc_feature
     70  <222> LOCATION: (1)...(697)
     71  <223> OTHER INFORMATION: n = A,T,C or G
     73  <400> SEQUENCE: 3
     74  gtactcagac ccccaacctc gaacagccag aagacaggtt gtctcctggg ccttggacac       60
W--> 75  agccngccag gccattgaag ganaagcaaa gacgaagcga accatctctc tccattgtgg      120
W--> 76  gggccaagta gctgcantan ccttcagtcc cagttgcatt gggttaaaga gctcatacat      180
W--> 77  actatgtgtn aggggtacag aagcttttcc tcatagggca tgagctctcc nagagttgac      240
W--> 78  cttttgcctn aacttggggt ttctgtggtt cataaagttn ggatatgtat ttttttttcaa     300
W--> 79  atggaanaaa atccgtattt ggcaaaaaga ctccaggggg atgatactgt ccttgccact      360
W--> 80  tacagtccaa angatnttcc ccaaagaata gacatttttt cctctcatca cttctggatg      420
W--> 81  caaaatcttt tatttttttc ctttctcgca ccnccccaga cccccttnnag gttnaaccgc      480
W--> 82  ttcccatctc ccccattcca cacgatnttg aattngcann ncgttgntgg tcgggtcccn      540
W--> 83  nccgaaaggg tnttttttatt cggggtnctg anttnnnaac cnctnagttg aatccgcggg      600
W--> 84  gcggccnngn gggttnnacc atgntgggga naactncccn ccgcgnttgg aatgccanag      660
W--> 85  ccttgaaant tttcttttgg tcgccccccn gagattc                             697
     87  <210> SEQ ID NO: 4
     88  <211> LENGTH: 712
     89  <212> TYPE: DNA
     90  <213> ORGANISM: Homo sapien
     92  <220> FEATURE:
     93  <221> NAME/KEY: misc_feature
     94  <222> LOCATION: (1)...(712)
     95  <223> OTHER INFORMATION: n = A,T,C or G
     97  <400> SEQUENCE: 4
W--> 98  gtactcagac aaccaatagg tgtgttyctc anatctgaaa cacaaaaaga ttctagctna       60
W--> 99  taatgttsaa tggttgaggg tttaagtgat cttggtatgt tngatttagc agcgatnggc      120
    100  cgggtgcggt ggctcacgca tgtatcccag cactttggga ggccgaggca ggaggatcac      180
W--> 101  ctgaggtcag gagtttgaga ccagcctggc cgacatggtn aaaccccgtc tctactanga      240
W--> 102  atacanaaat tagcccgggc atagtggcgc gtgcctrtga cctcsgctac tttggggatt      300
    103  ctcctgagga agaattgctt gaactcaggg aagtggargt ttgcagtgag cttaaatcaa      360
    104  gccactggca ctcccagcct gggktaacag agccamgact ctkgccgaaa aaaaraama      420
W--> 105  cgacggagaa nmagntctgt tattccatgg gaaattkgaa tttccttcyt tkaaatatct     480
W--> 106  taaaatnggt cctcctwaaa aaagttcggc tggggcccgk tggctcacat tttkttaycc     540
W--> 107  cyccccctttt tggggargggc caarggccgg kttgawtnnc ccttgagggg ccanaactcc    600
W--> 108  agnaaccrgn cccgggccar smgwkgkstr armccctttc cyyccmaraa aawwcsmaaa      660
    109  wwttycccsc cygsykggct ggkasckgtt myyyyygmtm csyagcttgc tt             712
    111  <210> SEQ ID NO: 5
    112  <211> LENGTH: 679
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Input Set : A:\475c7.app
Output Set: N:\CRF3\05132002\I667170A.raw

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     113 <212> TYPE: DNA
     114 <213> ORGANISM: Homo sapien
     116 <220> FEATURE:
     117 <221> NAME/KEY: misc_feature
     118 <222> LOCATION: (1)...(679)
     119 <223> OTHER INFORMATION: n = A,T,C or G
     121 <400> SEQUENCE: 5
W--> 122  gtactcagac cacctcacat gcagggtnag aaacatggag tgtgcggcag catcctcctc       60
     123  acatcccttt gtgagcacgg ctgctccgga atactgacca tctgggctag cacgacctaa      120
     124  cagagggttc tgcaggatgt gctattttaa agcagctggg tgcaacttgt gaaaacggga      180
W--> 125  atctngaagc agaacatgtn atcagcgatg gctgggattg gtggacagga ttgacaggag      240
     126  tatttgaggc tctaccaggc ctgtctacag gacagcttca tcgaagggac attttttaac      300
W--> 127  ctgttatttt anatnccaca tatntttttt aatgctnaag catacaggtt gaatttctgg      360
W--> 128  atcgtaacta ctagtgactt ctgaggttta cagttngaat atgttctcnn aggtttatca      420
W--> 129  agttntgtta ttgatgatng gtaatctaca cctctggaag ctgtngaatg tgaaaaagat      480
W--> 130  ncntncanct gaccagtttg nagggcactc tcttctggna agnaatccgn ccaaaaaaat      540
W--> 131  tgtttcnagg gggcntgggg ggtttaaaaa aatgtttctn ttnccntaaa aatgtttacc      600
W--> 132  cnnctattga aaaaatgggg gtcgnggggg gcttnaaatc cccnanttnt gaatnttnta      660
W--> 133  tccggaanct tggtttccc                                                   679
     135 <210> SEQ ID NO: 6
     136 <211> LENGTH: 369
     137 <212> TYPE: DNA
     138 <213> ORGANISM: Homo sapien
     140 <220> FEATURE:
     141 <221> NAME/KEY: misc_feature
     142 <222> LOCATION: (1)...(369)
     143 <223> OTHER INFORMATION: n = A,T,C or G
     145 <400> SEQUENCE: 6
     146  tcagtccagt catgggtcct ataagagaag tcactctgtg agtttccatg gaggaagaaa       60
     147  aagcttcatt tctttaccct gcagcaacag cggagggagg gagagcctat cttctttgca      120
W--> 148  aattcattaa ctttgtggtt gaaggagca gcgtcngaaa ctgctttagc acagtgggag      180
     149  gaaacaaac agattcatct ccggaaacca aaggaaaggg tragtgggtt tttattagcc      240
     150  agctgtatcc tagatggtca atttccagtg gatgaataca ccttacgtac gtttctcttg      300
W--> 151  cttcctacct nggcctgatc agctnggcac ttraatcatt ccgtnggggt wgctgtnaca      360
     152  ctggactga                                                             369
     154 <210> SEQ ID NO: 7
     155 <211> LENGTH: 264
     156 <212> TYPE: DNA
     157 <213> ORGANISM: Homo sapien
     159 <220> FEATURE:
     160 <221> NAME/KEY: misc_feature
     161 <222> LOCATION: (1)...(264)
     162 <223> OTHER INFORMATION: n = A,T,C or G
     164 <400> SEQUENCE: 7
W--> 165  tgctggatra gggatggggc acgggagcac agatmgactt taactgcccc cacgttntcm       60
     166  aggaaaggat tacaggcgtg agccactgcg cccggcctct tctccacttt cataggttcc      120
W--> 167  agtctctggt tcttctttct cagtttgttg tttttgcttc ttaaatmatg gagatnagaa      180
     168  tgaacactac actcggaatc aggaagccct gcctggcgcc tctgtcacct gtctaggggc      240
     169  ttcttctcac tgagtcatcc agca                                            264
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```
      171 <210> SEQ ID NO: 8
      172 <211> LENGTH: 280
      173 <212> TYPE: DNA
      174 <213> ORGANISM: Homo sapien
      176 <220> FEATURE:
      177 <221> NAME/KEY: misc_feature
      178 <222> LOCATION: (1)...(280)
      179 <223> OTHER INFORMATION: n = A,T,C or G
      181 <400> SEQUENCE: 8
W--> 182  acctcaactg cccanaacan aactgttgta caagatttga ggatttaaca atatttcaca     60
W--> 183  tgaaatattt cagacctacg ngagggctta aagacnaatt aaatgagcac cngtgtgccc    120
W--> 184  accgcccccna ttaagaatta gagcaagcag tgaggtgaag ccttgtcctt gcttttaaca   180
W--> 185  tagaaagtga tccaaattca ccaaacttga cttnnggttt tgcagtgtgg cctcctgatt     240
W--> 186  ctagacnctg gcgaaacatt tgatgggcaa aaaaaaaaa                            280
      188 <210> SEQ ID NO: 9
      189 <211> LENGTH: 449
      190 <212> TYPE: DNA
      191 <213> ORGANISM: Homo sapien
      193 <220> FEATURE:
      194 <221> NAME/KEY: misc_feature
      195 <222> LOCATION: (1)...(449)
      196 <223> OTHER INFORMATION: n = A,T,C or G
      198 <400> SEQUENCE: 9
W--> 199  tcgtcaactc caggatggct ttgaaaatna atggacacag atctctcctg ttttgatrat     60
W--> 200  ntgcagtgct natgactggc tttgcagttn attttgattc aggcaacaga tgttcctttt    120
      201  ggttccctgt ctcccatggg cgtcatttca tgttgtcctc tgccttcccc cagatattct    180
      202  aagttcagga cacaagcttc tggcccatgc agagcagagg ccatgagggg tcacagcatg    240
W--> 203  ggtacgggag gaaacactgg gctnacccag atnctggact tgagtcttgc ctctgctgct    300
      204  tgctgcacag cttctgtcat ggtgctaaac ctgtgacctg cctcacaggc ttagagcatg    360
W--> 205  cccgtagaag tactctnaac taaratgctt tccacaaatg agatggtttc atgaaaactt    420
      206  caaatagagg gcctgggcaa aaaaaaaaa                                      449
      208 <210> SEQ ID NO: 10
      209 <211> LENGTH: 538
      210 <212> TYPE: DNA
      211 <213> ORGANISM: Homo sapien
      213 <220> FEATURE:
      214 <221> NAME/KEY: misc_feature
      215 <222> LOCATION: (1)...(538)
      216 <223> OTHER INFORMATION: n = A,T,C or G
      218 <400> SEQUENCE: 10
      219  tttttttttt ttcccaaagg cctcaraaca ctagtcttct aattccaagc agaaagttac     60
      220  atccgccggg atacatgcca cttggtttga taaatcaaaa tacagcatcc ttcagatccc    120
W--> 221  tttgctgagc aatacaatta tttgtatatg ttactttttt ttctgtttgg ctnaaagatt    180
W--> 222  tgatatgagc tgaggaaaat gaagccntta ctgctatnag atctnatccc ttttccaccac   240
W--> 223  ctttcaggga tnttggcact gcayatattc agaattcccc nnagtcgctn gtgataaaaa    300
      224  tgtcttcaga gatggcagaa tatgtttcct ttggtacatg ttcattaaaa atatacacgt    360
W--> 225  gctcactact gtggatatgt atgtnttgac cgatnacaca ggctgattta gggaagagat    420
W--> 226  aaaagcacac ttngaattta ttagcctttc accnagacta anattctgaa attaagaatg    480
W--> 227  tattccttgg tcaacaattt tcctcttctc ttagccctct tacattgtan tggactga      538
```

RAW SEQUENCE LISTING
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Input Set : A:\475c7.app
Output Set: N:\CRF3\05132002\I667170A.raw

```
      229 <210> SEQ ID NO: 11
      230 <211> LENGTH: 543
      231 <212> TYPE: DNA
      232 <213> ORGANISM: Homo sapien
      234 <220> FEATURE:
      235 <221> NAME/KEY: misc_feature
      236 <222> LOCATION: (1)...(543)
      237 <223> OTHER INFORMATION: n = A,T,C or G
      239 <400> SEQUENCE: 11
      240   ttttttttt  ttgcccacag ctgccatctt tgtgtgataa ggccaacctt ctatgggaat      60
      241   caaccctcgc catcccagca aatcccctct ctcccttctc atgggagtgc cttgtattca     120
W--> 242   tcaggcatct gggacttgat gtgggtntgg gatttgaaat cagagcacct nggtctctst     180
W--> 243   caccattctn tcacttatta gctctnacct tgggtnaata cctgccttag tgtcntaggt     240
W--> 244   acaatatgaa tattgtctat ttctcaggga ttgcaatgac nagtnnatna gtgcatgaga     300
W--> 245   gggtaaaacc acagggtact ccgctcctcc naagaatgga gaattttttc tagaagccca     360
W--> 246   natntgcttg gaaggttggc caccnagagc cnnaatcttc ttttatttnc cactgaangc     420
W--> 247   ctaagaggna attctgaact catcccccnna tgacctctcc cgaatmagaa tatctctggc     480
W--> 248   acttaccata ttttcttgcc ctcttccact tacnaaactc ctttattcct taacnggacg     540
      249   aaa                                                                  543
      251 <210> SEQ ID NO: 12
      252 <211> LENGTH: 329
      253 <212> TYPE: DNA
      254 <213> ORGANISM: Homo sapien
      256 <400> SEQUENCE: 12
      257   cgatgacttg ggcagtgagt gggcctcctg ccaggtggca gggcacagct tagaccaaac      60
      258   ccttggcctc ccccctctgc agstacctct gaccaagaag gaaactagca agcctatgct     120
      259   ggcaagacca taggtggggt gctgggaatc ctcggggccg gctggcaccc actcctggtg     180
      260   ctcaagggag agacccactt gttcagatgc atggcctca ggcggttcaa ggcrgtctta     240
      261   gagccacaga gtcaaataaa aatcaatttt gagagaccac agcacctgct gctttgatcg     300
      262   tgatgttcaa ggcaagttgc aagtcatcg                                      329
      264 <210> SEQ ID NO: 13
      265 <211> LENGTH: 314
      266 <212> TYPE: DNA
      267 <213> ORGANISM: Homo sapien
      269 <220> FEATURE:
      270 <221> NAME/KEY: misc_feature
      271 <222> LOCATION: (1)...(314)
      272 <223> OTHER INFORMATION: n = A,T,C or G
      274 <400> SEQUENCE: 13
      275   cgatgacttg cacccgggag ctgtgacagt ggcctggaag cagatggcag ccccgtcaag      60
      276   gcgggagtgg agaccaccaa accctccaaa cagagcaaca actagtacgc ggccagcagc     120
      277   tacctgagcc tgacgcccga gcagtggaag tcccacagaa gctacagctg ccaggtcacg     180
W--> 278   catgaaggga gcaccgtgga gaagacagtg gccctacag aatgttcata ggttcccnac     240
W--> 279   tctnacccca cccacgggag cctgganctg cangatcccg ggggaagggt ctctctcccc     300
      280   atcccaagtc atcg                                                      314
      282 <210> SEQ ID NO: 14
      283 <211> LENGTH: 691
      284 <212> TYPE: DNA
      285 <213> ORGANISM: Homo sapien
```

```
RAW SEQUENCE LISTING ERROR SUMMARY      DATE: 05/13/2002
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```

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

```
Seq#:2;  N Pos. 79,459,463,466,469,585,642,645,656,660,666,670,671,678,686
Seq#:2;  N Pos. 693
Seq#:3;  N Pos. 65,83,137,140,190,231,250,280,307,372,376,453,467,468,474
Seq#:3;  N Pos. 507,515,519,520,521,527,540,541,552,567,572,575,576,577,582
Seq#:3;  N Pos. 585,607,608,610,616,617,624,631,636,640,646,658,669,690
Seq#:4;  N Pos. 32,59,102,117,220,238,246,431,435,487,578,579,594,603,610
Seq#:5;  N Pos. 28,185,200,312,315,324,337,396,409,410,425,439,465,481,483
Seq#:5;  N Pos. 485,488,501,519,523,530,547,555,580,583,586,602,603,625,635
Seq#:5;  N Pos. 644,646,649,655,658,668
Seq#:6;  N Pos. 156,311,325,345,357
Seq#:7;  N Pos. 57,176
Seq#:8;  N Pos. 15,20,81,96,112,129,214,215,247
Seq#:9;  N Pos. 29,61,71,90,264,273,377
Seq#:10; N Pos. 173,207,218,225,252,281,282,290,385,395,433,454,462,530
Seq#:11; N Pos. 147,171,190,206,216,235,281,285,286,289,331,361,364,385,392
Seq#:11; N Pos. 393,409,418,429,448,449,514,535
Seq#:13; N Pos. 238,244,267,273
Seq#:14; N Pos. 20,166,186,196,361,543,546,577,581,626,636,661
Seq#:15; N Pos. 87,146,195,333
Seq#:16; N Pos. 90,205,213,240,264,381,405,410,414,417,423,429,432,480,508
Seq#:233; N Pos. 90,97,242,244
Seq#:238; N Pos. 399
Seq#:245; N Pos. 105
Seq#:261; N Pos. 538
Seq#:264; N Pos. 32
Seq#:275; N Pos. 379
Seq#:279; N Pos. 457,471
Seq#:280; N Pos. 451
Seq#:281; N Pos. 456
Seq#:282; N Pos. 494
Seq#:290; N Pos. 233,247
Seq#:291; N Pos. 248,257,275,295,306,337
Seq#:292; N Pos. 19,353,415,431,474
Seq#:293; N Pos. 125,249,264,282,291,381,399,488
Seq#:294; N Pos. 55,285,489,519,582,590
Seq#:296; N Pos. 262,296,329,360,530,534,536,568,593,595
Seq#:298; N Pos. 105,108,136,148,149,237,243,250
Seq#:301; N Pos. 227,324,343,351,408,412,431,453,502,516,544
Seq#:302; N Pos. 51,157,240,258,304,316,378,391,475,576,580
Seq#:304; N Pos. 411
Seq#:306; N Pos. 502
Seq#:307; N Pos. 51,103,245,407,448
Seq#:308; N Pos. 576
Seq#:311; N Pos. 8,11,19
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Input Set : A:\475c7.app
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Seq#:313;  N Pos.  16,34,44,46,68,70,76,84,92,96,99,104,170,212,235,240,249
Seq#:313;  N Pos.  287,296,298,304,308,334,337,339,344,348,370,373
Seq#:314;  N Pos.  569
Seq#:315;  N Pos.  553
Seq#:318;  N Pos.  538,590
Seq#:320;  N Pos.  505,507,536,549,588
Seq#:321;  N Pos.  248,252
Seq#:322;  N Pos.  61,85,87,136,142,148,161,164,180,183,203,204,234,275,286
Seq#:322;  N Pos.  307,311,313,337,491,523,526
Seq#:323;  N Pos.  412,446
Seq#:324;  N Pos.  141,184,357
Seq#:325;  N Pos.  517
Seq#:326;  N Pos.  49,132
Seq#:328;  N Pos.  354
Seq#:329;  N Pos.  97,219,256,331,457
Seq#:330;  N Pos.  350,388,397,426,478,490,491
Seq#:331;  N Pos.  222,290,314,319,338,449,460
Seq#:334;  N Pos.  503,548
Seq#:336;  N Pos.  517
Seq#:337;  N Pos.  410
Seq#:338;  N Pos.  486,566,568
Seq#:339;  N Pos.  429
Seq#:340;  N Pos.  441,442
Seq#:342;  N Pos.  462,475
Seq#:344;  N Pos.  310,377
Seq#:346;  N Pos.  348,349
Seq#:353;  N Pos.  133,162,200,210,252,324,332,349,365,371
Seq#:354;  N Pos.  120,124,138,194,205,220,224,275,334
Seq#:355;  N Pos.  407,459,483
Seq#:356;  N Pos.  381
Seq#:357;  N Pos.  339,457,475,486
Seq#:358;  N Pos.  358,361,372,374,386
Seq#:359;  N Pos.  299,318,363,381,395,412
Seq#:360;  N Pos.  182,187,207,218,311,359,390,401,405,412,413,435,441
Seq#:361;  N Pos.  228,230,338,339,348,358,379
Seq#:362;  N Pos.  472,483,500,510
Seq#:369;  N Pos.  6,7
Seq#:370;  N Pos.  89,117,147,232
Seq#:371;  N Pos.  114,188,317
Seq#:372;  N Pos.  9
Seq#:376;  N Pos.  86,260
Seq#:381;  N Pos.  90,93
Seq#:389;  N Pos.  43,133,185,226,254,256,303
Seq#:399;  N Pos.  91,107,145,155,207,215,228,232,233,234,244
Seq#:401;  N Pos.  3
Seq#:406;  N Pos.  402,414
Seq#:409;  N Pos.  279,304,315,319,343,352,355,356,361,381
Seq#:411;  N Pos.  370
Seq#:415;  N Pos.  380,394
```

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 05/13/2002
PATENT APPLICATION: **US/09/667,170A**   TIME: 15:12:13

Input Set : **A:\475c7.app**
Output Set: **N:\CRF3\05132002\I667170A.raw**

Seq#:416; N Pos. 180,202,206,214,224,235,240,241,245,247,250,251,252,257
Seq#:416; N Pos. 267

VERIFICATION SUMMARY   DATE: 05/13/2002
PATENT APPLICATION: US/09/667,170A   TIME: 15:12:13

Input Set : A:\475c7.app
Output Set: N:\CRF3\05132002\I667170A.raw

L:20 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:51 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:60
L:57 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:420
L:59 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:540
L:60 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:600
L:61 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:660
L:75 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:60
L:76 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:120
L:77 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:180
L:78 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:240
L:79 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:300
L:80 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:360
L:81 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:420
L:82 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:480
L:83 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:540
L:84 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:600
L:85 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:660
L:98 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
L:99 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:60
L:101 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:180
L:102 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:240
L:105 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:420
L:106 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:480
L:107 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:540
L:108 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:600
L:122 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
L:125 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:180
L:127 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:300
L:128 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:360
L:129 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:420
L:130 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:480
L:131 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:540
L:132 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:600
L:133 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:660
L:148 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:120
L:151 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:300
L:165 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:167 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:120
L:182 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:183 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:60
L:184 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:120
L:185 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:180
L:186 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:240
L:199 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:200 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:60
L:203 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:240
L:205 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:360
L:221 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:120

VERIFICATION SUMMARY   DATE: 05/13/2002
PATENT APPLICATION: US/09/667,170A   TIME: 15:12:13

Input Set : A:\475c7.app
Output Set: N:\CRF3\05132002\I667170A.raw

L:222 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:180
L:223 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:240
L:225 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:360
L:6769 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:279